

In the Specification

Please replace the table on page 52 of the specification with the following table:

	/DB_XREF=gi:7656972 /UG=Hs.241535 apolipoprotein L, 3 /FL=gb:AY014905.1 gb:AF070675.1 gb:NM_014349.1
	gb:NM_018229.1 /DEF=Homo sapiens hypothetical protein FLJ10813 (FLJ10813), mRNA. /FEA=mrna /GEN=FLJ10813
218139 s at	/PROD=hypothetical protein FLJ10813 /DB_XREF=gi:8922687 /UG=Hs.106210 hypothetical protein FLJ10813
	/FL=gb:AL136685.1 gb:NM_018229.1
213878 at	Consensus includes gb:AI685944 /FEA=EST /DB_XREF=gi:4897238 /DB_XREF=est:tu38g02.x1 /CLONE=IMAGE:2253362
	/UG=Hs.235069 RecQ protein-like (DNA helicase Q1-like)
	gb:NM_024551.1 /DEF=Homo sapiens hypothetical protein FLJ21432 (FLJ21432), mRNA. /FEA=mrna /GEN=FLJ21432
	/PROD=hypothetical protein FLJ21432 /DB_XREF=gi:13375714 /UG=Hs.11641 hypothetical protein FLJ21432
201346 at	/FL=gb:NM_024551.1 gb:BC004906.1
	Consensus includes gb:AL138761 /DEF=Human DNA sequence from clone RP11-16H23 on chromosome 10.
	Contains the gene KIAA0204 (HSLK) for a protein kinase, the COL17A1 gene for collagen type XVII alpha 1 (BP180),
	ESTs and GSSs /FEA=mrna 2 /DB_XREF=gi:8573811 /UG=Hs.105751 Ste20-related serinethreonine kinase
206874 s at	/FL=gb:DB6959.1 gb:NM_014720.1
	Consensus includes gb:NM_004251.1 /DEF=Homo sapiens RAB9, member RAS oncogene family (RAB9), mRNA. /FEA=CDS /GEN=RAB9
221808 at	/PROD=RAB9, member RAS oncogene family /DB_XREF=gi:4759011 /UG=Hs.28726 RAB9, member RAS oncogene family
	/FL=gb:U44103.1 gb:NM_004251.1
	gb:BC005297.1 /DEF=Homo sapiens, Similar to kynurenine 3-monooxygenase (kynurenine 3-hydroxylase), clone MGC:12362, mRNA,
	complete cds. /FEA=mrna /PROD=Similar to kynurenine 3-monooxygenase(kynurenine 3-hydroxylase) /DB_XREF=gi:13529016
211138 s at	/UG=Hs.107318 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) /FL=gb:BC005297.1
	gb:AF279891.1 /DEF=Homo sapiens dead box protein 15 mRNA, complete cds. /FEA=mrna /PROD=dead box protein 15
	/DB_XREF=gi:19624452 /UG=Hs.5683 DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 15 /FL=gb:AB001636.1
201386 s at	gb:NM_001358.1 gb:AF279891.1
	gb:NM_014751.1 /DEF=Homo sapiens KIAA0429 gene product (KIAA0429), mRNA. /FEA=mrna /GEN=KIAA0429 /PROD=KIAA0429 gene
	product
203037 s at	/DB_XREF=gi:7662113 /UG=Hs.77694 KIAA0429 gene product /FL=gb:AB007889.1 gb:NM_014751.1
	gb:NM_013393.1 /DEF=Homo sapiens cell division protein FtsJ (FtsJ), mRNA. /FEA=mrna /GEN=FUH1 /PROD=cell division protein
218356 at	FtsJ /DB_XREF=gi:7019376 /UG=Hs.279877 cell division protein FtsJ /FL=gb:AF093415.1 gb:NM_013393.1
	gb:NM_014473.1 /DEF=Homo sapiens putative dimethyladenosine transferase (HSA9761), mRNA. /FEA=mrna /GEN=HSA9761
	/PROD=putative dimethyladenosine transferase /DB_XREF=gi:7657197 /UG=Hs.125819 putative dimethyladenosine transferase
204405 x at	/FL=gb:AF102147.1 gb:NM_014473.1
	gb:NM_007062.1 /DEF=Homo sapiens nuclear phosphoprotein similar to S. cerevisiae PWP1 (PWP1), mRNA. /FEA=mrna /GEN=PWPI
	/PROD=nuclear phosphoprotein similar to S. cerevisiaePWP1 /DB_XREF=gi:5902033
201608 s at	/UG=Hs.172589 nuclear phosphoprotein similar to S. cerevisiae_PWP1 /FL=gb:BC001652.1 gb:L07758.1 gb:NM_007062.1
	gb:NM_015380.1 /DEF=Homo sapiens CGI-51 protein (CGI-51), mRNA. /FEA=mrna /GEN=CGI-51 /PROD=CGI-51 protein
201569 s at	/DB_XREF=gi:7661541
	/UG=Hs.4877 CGI-51 protein /FL=gb:AF151809.1 gb:NM_015380.1
222244 s at	Consensus includes gb:AK000749.1 /DEF=Homo sapiens CDNA FLJ20742 fis, clone HEP06891. /FEA=mrna /DB_XREF=gi:7021031
	/UG=Hs.52184 hypothetical protein FLJ20618
	gb:NM_004939.1 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 1 (DDX1), mRNA. /FEA=mrna
	/GEN=DDX1 /PROD=DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 1 /DB_XREF=gi:4826685 /UG=Hs.78580 DEADH (Asp-Glu-
201241 at	Ala-AspHis) (SEQ ID NO:1) box polypeptide 1 /FL=gb:X70649.1 gb:NM_004939.1
	gb:NM_018090.1 /DEF=Homo sapiens hypothetical protein FLJ10420 (FLJ10420), mRNA. /FEA=mrna /GEN=FLJ10420
220731 s at	/PROD=hypothetical protein FLJ10420 /DB_XREF=gi:8922415 /UG=Hs.289087 hypothetical protein FLJ10420 /FL=gb:NM_018090.1
208799 at	gb:BC004146.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, beta type, 5, clone MGC:2175, mRNA, complete
	cds.

Please replace the table on page 64 of the specification with the following table:

206715_at	gb:NM_012252.1 /DEF=Homo sapiens transcription factor EC (TFEC), mRNA. /FEA=mrna /GEN=TFEC /PROD=transcription factor EC /DB_XREF=gi:6912701 /UG=Hs.113274 transcription factor EC /FL=gb:AB014580.1 gb:NM_012252.1
213123_at	Consensus includes gb:BE222709 /FEA=EST /DB_XREF=gi:8910027 /DB_XREF=est:hu51g06.x1 /CLONE=IMAGE:3173626 /UG=Hs.28785 microfibrillar-associated protein 3
204049_s_at	gb:NM_014721.1 /DEF=Homo sapiens KIAA0680 gene product (KIAA0680), mRNA. /FEA=mrna /GEN=KIAA0680 /PROD=KIAA0680 gene product /DB_XREF=gi:7662247 /UG=Hs.102471 KIAA0680 gene product /FL=gb:AB014580.1 gb:NM_014721.1
201985_at	gb:NM_014846.1 /DEF=Homo sapiens KIAA0196 gene product (KIAA0196), mRNA. /FEA=mrna /GEN=KIAA0196 /PROD=KIAA0196 gene product /DB_XREF=gi:7661987 /UG=Hs.8294 KIAA0196 gene product /FL=gb:DB3780.1 gb:NM_014846.1
208773_s_at	gb:AL136943.1 /DEF=Homo sapiens mRNA; cDNA DKF2p586G1024 (from clone DKF2p586G1024); complete cds. /FEA=mrna /GEN=DKF2p586G1024 /PROD=hypothetical protein /DB_XREF=gi:12053380 /UG=Hs.301226 KIAA1085 protein /FL=gb:AL136943.1
219563_at	gb:NM_024633.1 /DEF=Homo sapiens hypothetical protein FLJ21276 (FLJ21276), mRNA. /FEA=mrna /GEN=FLJ21276 /PROD=hypothetical protein FLJ21276 /DB_XREF=gi:13375863 /UG=Hs.41502 hypothetical protein FLJ21276 /FL=gb:NM_024633.1
218501_at	gb:NM_019555.1 /DEF=Homo sapiens Rho guanine nucleotide exchange factor (GEF) 3 (ARHGGEF3), mRNA. /FEA=mrna /GEN=ARHGGEF3 /PROD=Rho guanine nucleotide exchange factor (GEF) 3 /DB_XREF=gi:9506400 /UG=Hs.25951 Rho guanine nucleotide exchange factor (GEF) 3 /FL=gb:AF249744.1 gb:NM_019555.1
212833_at	Consensus includes gb:M74089.1 /DEF=Human TBI gene mRNA, 3 end. /FEA=mrna /GEN=TBI /DB_XREF=gi:182400 /UG=Hs.75639 Human TBI gene mRNA, 3 end
209623_at	Consensus includes gb:AW439494 /FEA=EST /DB_XREF=gi:6974800 /DB_XREF=est:xtl9c01.x1 /CLONE=IMAGE:2779584 /UG=Hs.167531 methylcrotonoyl-Coenzyme A carboxylase 2 (beta) /FL=gb:AB050049.1 gb:AF310971.1 gb:AF301000.1 gb:NM_022132.2
209969_s_at	gb:BC002704.1 /DEF=Homo sapiens, Similar to signal transducer and activator of transcription 1, 91kD, clone MGC:3493, mRNA, complete cds. /FEA=mrna /PROD=Similar to signal transducer and activator of transcription 1, 91kD /DB_XREF=gi:12803734 /UG=Hs.21486 signal transducer and activator of transcription 1, 91kD /FL=gb:BC002704.1
219966_x_at	gb:NM_017869.1 /DEF=Homo sapiens BANP homolog, SMAR1 homolog (FLJ20538), mRNA. /FEA=mrna /GEN=FLJ20538 /PROD=BANP homolog, SMAR1 homolog /DB_XREF=gi:8923506 /UG=Hs.194637 BANP homolog, SMAR1 homolog /FL=gb:NM_017869.1
213275_x_at	Consensus includes gb:W47179 /FEA=EST /DB_XREF=gi:1332046 /DB_XREF=est:zc34d07.s1 /CLONE=IMAGE:324205 /UG=Hs.297939 cathepsin B
210231_x_at	gb:D45198.1 /DEF=Human mRNA for template activating factor-I alpha, complete cds. /FEA=mrna /GEN=template activating factor-I alpha /DB_XREF=gi:971271 /UG=Hs.145279 SET translocation (myeloid leukemia-associated) /FL=gb:D45198.1
212474_at	Consensus includes gb:D87682.1 /DEF=Human mRNA for KIAA0241 gene, partial cds. /FEA=mrna /GEN=KIAA0241 /DB_XREF=gi:1663699 /UG=Hs.150275 KIAA0241 protein
208717_at	gb:BC001669.1 /DEF=Homo sapiens, Similar to oxidase (cytochrome c) assembly 1-like, clone MGC:2171, mRNA, complete cds. /FEA=mrna /PROD=Similar to oxidase (cytochrome c) assembly 1-like /DB_XREF=gi:12804516 /UG=Hs.151134 oxidase (cytochrome c) assembly 1-like /FL=gb:BC001669.1 gb:NM_005015.1
217527_s_at	Consensus includes gb:AI478300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALU6 HUMAN ALU SUBFAMILY SP SEQUENCE CONTAMINATION WARNING ENTRY H.sapiens
220495_s_at	gb:NM_024715.1 /DEF=Homo sapiens hypothetical protein FLJ22625 (FLJ22625), mRNA. /FEA=mrna /GEN=FLJ22625 /PROD=hypothetical protein FLJ22625 /DB_XREF=gi:13376016 /UG=Hs.106534 hypothetical protein FLJ22625 /FL=gb:NM_024715.1
200892_s_at	gb:BC000451.1 /DEF=Homo sapiens, splicing factor, arginineserine-rich (transformer 2 Drosophila homolog) 10, clone MGC:8454, mRNA, complete cds. /FEA=mrna /PROD=splicing factor, arginineserine-rich (transformer 2 Drosophila homolog) 10 /DB_XREF=gi:12653362 /UG=Hs.30035 splicing factor, arginineserine-rich (transformer 2 Drosophila homolog) 10 /FL=gb:BC000451.1 gb:BC000451.1 gb:U61267.1 gb:U68063.1 gb:NM_004593.1
201798_s_at	gb:NM_013451.1 /DEF=Homo sapiens fer-1 (C.elegans)-like 3 (myoferlin) (FER1L3), mRNA. /FEA=mrna /GEN=FER1L3 /PROD=fer-1 (C.elegans)-like 3 (myoferlin) /DB_XREF=gi:7305052 /UG=Hs.234680 fer-1 (C.elegans)-like 3 (myoferlin) /FL=gb:AF182316.1 gb:NM_013451.1
202529_at	gb:NM_002766.1 /DEF=Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 1 (PRPSAP1), mRNA. /FEA=mrna /GEN=PRPSAP1 /PROD=phosphoribosyl pyrophosphatesynthetase-associated protein 1 /DB_XREF=gi:4506130 /UG=Hs.77498 phosphoribosyl pyrophosphate synthetase-associated protein 1 /FL=gb:D61391.1 gb:NM_002766.1
208897_s_at	gb:BC003360.1 /DEF=Homo sapiens, DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 (Myc-regulated), clone MGC:5316, mRNA, complete cds. /FEA=mrna /PROD=DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 (Myc-regulated) /DB_XREF=gi:13097182 /UG=Hs.100555 DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 (Myc-regulated) /FL=gb:BC001238.1 gb:BC003360.1
210296_s_at	gb:BC005375.1 /DEF=Homo sapiens, peroxisomal membrane protein 3 (35kD, Zellweger syndrome), clone MGC:12491, mRNA, complete cds. /FEA=mrna /PROD=peroxisomal membrane protein 3 (35kD, Zellwegersyndrome) /DB_XREF=gi:13529226 /UG=Hs.180612 peroxisomal membrane protein 3 (35kD, Zellweger syndrome) /FL=gb:BC005375.1
212460_at	Consensus includes gb:BE738425 /FEA=EST /DB_XREF=gi:10152417 /DB_XREF=est:601572441T1 /CLONE=IMAGE:3839147 /UG=Hs.241507 ribosomal protein S6

Please replace the table on page 66 of the specification with the following table:

	protein /FL=gb:AF178930.1 gb:NM_022162.1
218583_s_at	gb:NM_020640.1 /DEF=Homo sapiens RP42 homolog (RP42), mRNA. /FEA=mRNA /GEN=RP42 /PROD=RP42 homolog /DB_XREF=gi:10190677 /UG=Hs.104613 RP42 homolog /FL=gb:NM_020640.1 gb:AF292100.2
221689_s_at	gb:AB035745.1 /DEF=Homo sapiens mRNA for DSCR5b, complete cds. /FEA=mRNA /GEN=DSCR5b /PROD=DSCR5b /DB_XREF=gi:7798596 /UG=Hs.66493 Down syndrome critical region gene 5 /FL=gb:AB035745.1 gb:AB037163.1 gb:AF237812.1
205412_at	gb:NM_000019.1 /DEF=Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ACAT1 /PROD=acetyl-Coenzyme A acetyltransferase 1 precursor /DB_XREF=gi:4557236 /UG=Hs.37 acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase) /FL=gb:NM_000019.1
202542_s_at	gb:NM_004757.1 /DEF=Homo sapiens small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating) (SCYE1), mRNA. /FEA=mRNA /GEN=SCYE1 /PROD=small inducible cytokine subfamily E, member 1 /DB_XREF=gi:4758265 /UG=Hs.146401 small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating) /FL=gb:NM_004757.1 gb:U10117.1
202521_at	gb:NM_006565.1 /DEF=Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA. /FEA=mRNA /GEN=CTCF /PROD=CCCTC-binding factor (zinc finger protein) /DB_XREF=gi:5729789 /UG=Hs.57419 CCCTC-binding factor (zinc finger protein) /FL=gb:NM_006565.1 gb:U25435.1
204391_x_at	gb:NM_015905.1 /DEF=Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA. /FEA=mRNA /GEN=TIF1 /PROD=transcriptional intermediary factor 1 alpha /DB_XREF=gi:7706233 /UG=Hs.183858 transcriptional intermediary factor 1 /FL=gb:AF009353.1 gb:AF119042.1 gb:NM_003852.1 gb:NM_015905.1
214733_s_at	Consensus includes gb:AL031427 /DEF=Human DNA sequence from clone 167A19 on chromosome 1p32.1-33. Contains three genes for novel proteins, the DIO1 gene for type I iodothyronine deiodinase (EC 3.8.1.4, TXDI1, ITDI1) and an HNRNP A3 (Heterogenous Nuclear Ribonucleoprotein A3, FBRNP) ... /FEA=mRNA 6 /DB_XREF=gi:4835258 /UG=Hs.11923 hypothetical protein
217864_s_at	gb:NM_016166.1 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box binding protein 1 (DDXBPL), mRNA. /FEA=mRNA /GEN=DDXBPL /PROD=DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box binding protein 1 /DB_XREF=gi:7706636 /UG=Hs.75251 DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box binding protein 1 /FL=gb:AF077951.1 gb:AF167160.1 gb:NM_016166.1
212904_at	Consensus includes gb:AB033011.1 /DEF=Homo sapiens mRNA for KIAA1185 protein, partial cds. /FEA=mRNA /GEN=KIAA1185 /PROD=KIAA1185 protein /DB_XREF=gi:6330301 /UG=Hs.268488 KIAA1185 protein
202126_at	Consensus includes gb:AA156948 /FEA=EST /DB_XREF=gi:1728563 /DB_XREF=est:z119f02.s1 /CLONE=IMAGE:502395 /UG=Hs.198891 serinethreonine-protein kinase PRP4 homolog /FL=gb:U48736.1 gb:NM_003913.1
203484_at	gb:NM_014302.1 /DEF=Homo sapiens Sec61 gamma (SEC61G), mRNA. /FEA=mRNA /GEN=SEC61G /PROD=Sec61 gamma /DB_XREF=gi:7657545 /UG=Hs.9950 Sec61 gamma /FL=gb:AF054184.1 gb:NM_014302.1
203345_s_at	Consensus includes gb:AI566096 /FEA=EST /DB_XREF=gi:4524548 /DB_XREF=est:tn53d02.x1 /CLONE=IMAGE:21172099 /UG=Hs.31016 putative DNA binding protein /FL=gb:AF072814.1 gb:NM_007358.1
213238_at	Consensus includes gb:AI478147 /FEA=EST /DB_XREF=gi:4371373 /DB_XREF=est:tm34f06.x1 /CLONE=IMAGE:2160035 /UG=Hs.173540 ATPase, Class V, type 10D
202680_at	gb:NM_002095.1 /DEF=Homo sapiens general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) (GTF2E2), mRNA. /FEA=mRNA /GEN=GTF2E2 /PROD=general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) /DB_XREF=gi:4504194 /UG=Hs.77100 general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) /FL=gb:NM_002095.1
218117_at	gb:NM_014248.1 /DEF=Homo sapiens ring-box 1 (RBX1), mRNA. /FEA=mRNA /GEN=RBX1 /PROD=ring-box 1 /DB_XREF=gi:7657507 /UG=Hs.279919 ring-box 1 /FL=gb:BC001466.1 gb:AF140598.1 gb:AF142059.1 gb:NM_014248.1
218768_at	gb:NM_020401.1 /DEF=Homo sapiens nuclear pore complex protein (NUP107), mRNA. /FEA=mRNA /GEN=NUP107 /PROD=nuclear pore complex protein /DB_XREF=gi:9966880 /UG=Hs.236204 nuclear pore complex protein /FL=gb:NM_020401.1
202271_at	Consensus includes gb:AB007952.1 /DEF=Homo sapiens mRNA for KIAA0483 protein, partial cds. /FEA=mRNA /GEN=KIAA0483 /PROD=KIAA0483 protein /DB_XREF=gi:3413925 /UG=Hs.64691 KIAA0483 protein /FL=gb:NM_015176.1
218543_s_at	gb:NM_022750.1 /DEF=Homo sapiens hypothetical protein FLJ22693 (FLJ22693), mRNA. /FEA=mRNA /GEN=FLJ22693 /PROD=hypothetical protein FLJ22693 /DB_XREF=gi:12232412 /UG=Hs.12646 hypothetical protein FLJ22693 /FL=gb:AL136766.1 gb:NM_022750.1
203146_s_at	gb:NM_001470.1 /DEF=Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1), transcript variant 1, mRNA. /FEA=mRNA /GEN=GABBR1 /PROD=gamma-aminobutyric acid (GABA) B receptor 1, isoform A precursor /DB_XREF=gi:10835014 /UG=Hs.167017 gamma-aminobutyric acid (GABA) B receptor, 1 /FL=gb:NM_001470.1 gb:AF301005.1 gb:AF099148.1
218140_x_at	gb:NM_021203.1 /DEF=Homo sapiens APMCF1 protein (APMCF1), mRNA. /FEA=mRNA /GEN=APMCF1 /PROD=APMCF1 protein /DB_XREF=gi:10864014 /UG=Hs.12152 APMCF1 protein /FL=gb:NM_021203.1 gb:AF141882.1
40420_at	Cluster Incl. AB015718:Homo sapiens lok mRNA for protein kinase, complete cds /cds=(50,2956) /gb=AB015718 /gi=4001687 /ug=Hs.16134 /len=4221

Please replace the table on page 77 of the specification with the following table:

202399_s_at	59kd /FL=gb:AL136798.1 gb:BC000480.1 gb:U50939.1 gb:NM_003905.1
218515_at	gb:NM_005829.1 /DEF=Homo sapiens adaptor-related protein complex 3, sigma 2 subunit (AP3S2), mRNA. /FEA=mRNA /GEN=AP3S2 /PROD=adaptor-related protein complex 3, sigma 2 subunit /DB_XREF=gi:5031580 /UG=Hs.154782 adaptor-related protein complex 3, sigma 2 subunit /FL=gb:BC002785.1 gb:NM_005829.1
209180_at	gb:NM_016631.1 /DEF=Homo sapiens hypothetical protein (LOC51325), mRNA. /FEA=mRNA /GEN=LOC51325 /PROD=hypothetical protein /DB_XREF=gi:7706175 /UG=Hs.26461 hypothetical protein /FL=gb:AF208862.1 gb:NM_016631.1
205105_at	gb:U49245.1 /DEF=Human geranylgeranyl transferase type II beta-subunit mRNA, complete cds. /FEA=mRNA /PROD=geranylgeranyl transferase type II beta-subunit /DB_XREF=gi:1216503 /UG=Hs.78948 Rab geranylgeranyltransferase, beta subunit /FL=gb:U49245.1 gb:NM_004582.1
200666_s_at	gb:NM_002372.1 /DEF=Homo sapiens mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA. /FEA=mRNA /GEN=MAN2A1 /PROD=mannosidase, alpha, class 2A, member 1 /DB_XREF=gi:4758697 /UG=Hs.32965 mannosidase, alpha, class 2A, member 1 /FL=gb:U31520.1 gb:NM_002372.1 gb:D63998.1
200972_at	gb:NM_006145.1 /DEF=Homo sapiens heat shock 40kd protein 1 (HSPF1), mRNA. /FEA=mRNA /GEN=HSPF1 /PROD=heat shock 40kd protein 1 /DB_XREF=gi:5453689 /UG=Hs.82646 DnaJ (Hsp40) homolog, subfamily B, member 1 /FL=gb:BC002352.1 gb:NM_006145.1 gb:D49547.1
218352_at	gb:BC000704.1 /DEF=Homo sapiens, tetraspan 3, clone MGC:965, mRNA, complete cds. /FEA=mRNA /PROD=tetraspan 3 /DB_XREF=gi:12653830 /UG=Hs.100090 tetraspan 3 /FL=gb:BC000704.1 gb:AF054840.1 gb:NM_005724.1 gb:AF133423.1
217728_at	gb:NM_018191.1 /DEF=Homo sapiens hypothetical protein FLJ10716 (FLJ10716), mRNA. /FEA=mRNA /GEN=FLJ10716 /PROD=hypothetical protein FLJ10716 /DB_XREF=gi:8922616 /UG=Hs.24129 hypothetical protein FLJ10716 /FL=gb:NM_018191.1
211971_s_at	gb:NM_014624.2 /DEF=Homo sapiens S100 calcium-binding protein A6 (calyculin) (S100A6), mRNA. /FEA=mRNA /GEN=S100A6 /PROD=S100 calcium-binding protein A6 /DB_XREF=gi:9845517 /UG=Hs.275243 S100 calcium-binding protein A6 (calyculin) /FL=gb:BC001431.1 gb:NM_014624.2
212500_at	Consensus includes gb:AI653608 /FEA=EST /DB_XREF=gi:4737587 /DB_XREF=est:tz21a06.x1 /CLONE=IMAGE:2289202 /UG=Hs.182490 leucine-rich protein mRNA
218473_s_at	Consensus includes gb:AL049319.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C046 (from clone DKFZp564C046). /FEA=mRNA /DB_XREF=gi:4500092 /UG=Hs.99821 Homo sapiens mRNA; cDNA DKFZp564C046 (from clone DKFZp564C046)
203580_s_at	gb:NM_024656.1 /DEF=Homo sapiens hypothetical protein FLJ22329 (FLJ22329), mRNA. /FEA=mRNA /GEN=FLJ22329 /PROD=hypothetical protein FLJ22329 /DB_XREF=gi:13375904 /UG=Hs.61478 hypothetical protein FLJ22329 /FL=gb:NM_024656.1
200900_s_at	gb:NM_003983.1 /DEF=Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 (SLC7A6), mRNA. /FEA=mRNA /GEN=SLC7A6 /PROD=solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 /DB_XREF=gi:4507052 /UG=Hs.10315 solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 /FL=gb:D87432.1 gb:NM_003983.1
221652_s_at	Consensus includes gb:AI583537 /FEA=EST /DB_XREF=gi:4569434 /DB_XREF=est:ts12d03.x1 /CLONE=IMAGE:2228357 /UG=Hs.75709 mannose-6-phosphate receptor (cation dependent) /FL=gb:NM_002355.2 gb:MI6985.1
217750_s_at	gb:AF274950.1 /DEF=Homo sapiens PNAS-25 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-25 /DB_XREF=gi:12751064 /UG=Hs.22595 hypothetical protein FLJ10637 /FL=gb:AF274950.1
203544_s_at	gb:NM_023079.1 /DEF=Homo sapiens hypothetical protein FLJ13855 (FLJ13855), mRNA. /FEA=mRNA /GEN=FLJ13855 /PROD=hypothetical protein FLJ13855 /DB_XREF=gi:12751494 /UG=Hs.168232 hypothetical protein FLJ13855 /FL=gb:NM_023079.1
221580_s_at	gb:NM_003473.1 /DEF=Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (STAM), mRNA. /FEA=mRNA /GEN=STAM /PROD=signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /DB_XREF=gi:4507248 /UG=Hs.153487 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /FL=gb:U43899.1 gb:NM_003473.1
202629_at	gb:BC001972.1 /DEF=Homo sapiens, clone MGC:5306, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:5306) /DB_XREF=gi:12805036 /UG=Hs.301732 hypothetical protein MGC5306 /FL=gb:BC001972.1
205763_s_at	Consensus includes gb:AV681579 /FEA=EST /DB_XREF=gi:10283442 /DB_XREF=est:AV681579 /CLONE=GKBAFE05 /UG=Hs.84084 amyloid beta precursor protein (cytoplasmic tail)-binding protein 2 /FL=gb:AF01782.1 gb:NM_006380.1
212337_at	gb:NM_006773.2 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 (Myc-regulated) (DDX18), mRNA. /FEA=mRNA /GEN=DDX18 /PROD=DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 /DB_XREF=gi:13787205 /UG=Hs.100555 DEADH (Asp-Glu-Ala-AspHis) (SEQ ID NO:1) box polypeptide 18 (Myc-regulated) /FL=gb:NM_006773.2
212244_at	Consensus includes gb:AI687738 /FEA=EST /DB_XREF=gi:4899032 /DB_XREF=est:tp93g08.x1 /CLONE=IMAGE:2206910 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin B)
222011_s_at	Consensus includes gb:AL050091.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1918 (from clone DKFZp586F1918); partial cds. /FEA=mRNA /GEN=DKFZp586F1918 /PROD=hypothetical protein /DB_XREF=gi:4884111 /UG=Hs.6283 DKFZp586F1918 protein
204725_s_at	Consensus includes gb:BF224073 /FEA=EST /DB_XREF=gi:11131299 /DB_XREF=est:7q83e05.x1 /CLONE=IMAGE:3704936 /UG=Hs.278544 acetyl-Coenzyme A acetyltransferase 2 (acetoacetyl Coenzyme A thiolase)
	gb:NM_006153.1 /DEF=Homo sapiens NCK adaptor protein 1 (NCK1), mRNA. /FEA=mRNA /GEN=NCK1 /PROD=NCK adaptor protein 1 /DB_XREF=gi:5453753 /UG=Hs.54589 NCK adaptor protein 1 /FL=gb:NM_006153.1

Please insert the accompanying Sequence Listing as new page 1 following page 96 (Abstract of the Disclosure) in the subject specification.